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Original Article

Fabrication and Evaluation of PVA-NYS-THY Nanofiber Scaffolds as Antifungal Agents Against Fluconazole-resistant *Candida glabrata*

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ABSTRACT

Introduction: The emergence of fluconazole-resistant *Candida glabrata* presents a significant challenge in antifungal therapy, necessitating the development of alternative treatment strategies. *C. glabrata*, an opportunistic yeast, is increasing resistance to common antifungals like fluconazole, often due to efflux pump overexpression, leading to compromised treatment efficacy, higher mortality, prolonged hospital stays, and increased healthcare costs. This study focused on fabricating and evaluating polyvinyl alcohol-nystatin-thymol (PVA-NYS-THY) nanofibrous scaffolds as a novel antifungal approach against fluconazole-resistant *C. glabrata*.

Materials & Methods: Clinical isolates were identified and assessed for resistance using culture methods, molecular assays, and antifungal susceptibility testing. PVA-NYS-THY nanofibers, produced via electrospinning, exhibited uniform fibers with an average diameter of ~100 nm as confirmed by scanning electron microscopy (SEM). Fourier transform infrared spectroscopy confirmed the successful incorporation of functional groups. Real-time polymerase chain reaction (PCR) was employed to evaluate the nanofibers' effect on the expression of secreted aspartyl proteinases (*SAP*) and agglutinin-like sequence (*ALS*) gene. Scaffold release kinetics were characterized, and antifungal efficacy was assessed using minimum inhibitory concentration (MIC) assays.

Results: PVA-NYS-THY scaffolds showed favorable release profiles and significantly downregulated *ALS* and *SAP* gene expression. MIC values for PVA-NYS-THY, PVA-NYS, and PVA-THY were 7.81, 15.62, and 62.5 µg/mL, respectively, demonstrating superior antifungal activity of the PVA-NYS-THY formulation.

Conclusion: These findings suggest PVA-NYS-THY nanofibrous scaffolds offer a promising therapeutic strategy for combating fluconazole-resistant *C. glabrata*, providing a novel solution to overcome current limitations in antifungal treatment.

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1. Introduction

Candida glabrata, an opportunistic yeast within the *Candida* genus, presents a growing threat to human health, particularly due to its increasing antifungal resistance to antifungal agents. Candidiasis, the infection caused by *Candida* species, can affect various anatomical locations, including the oral cavity, esophagus, vagina, urinary tract, skin, and bloodstream [1]. Immunocompromised individuals are particularly susceptible to severe candidiasis [2]. Effective management of these infections is challenged by availability of antifungal drugs and the rise of drug resistance, especially in *C. glabrata*, which frequently exhibits azole resistance due to efflux pump overexpression or *ERG11* gene mutations [3]. This resistance leads to high mortality, prolonged hospital stays, and increased healthcare costs [4], necessitating innovative therapeutic strategies. Several approaches have been explored to combat fungal infections, including conventional antifungal drugs (e.g. azoles, polyenes, echinocandins) and various drug delivery systems. Traditional antifungal therapies often suffer from limitations such as poor solubility, instability, limited bioavailability, and systemic toxicity [5]. For instance, topical formulations have limited penetration into deeper tissues, while oral and intravenous administrations can cause adverse side effects. Furthermore, the emergence of drug resistance significantly reduces the efficacy of these conventional treatments. This highlights the need for improved drug delivery systems.

Nanomaterials, particularly nanofibers, offer a promising alternative by addressing these limitations. These ultrafine fibers (diameter <1000 nm) possess a high surface area-to-volume ratio, tunable porosity, flexibility, and mechanical strength, all of which contribute to enhanced drug delivery [5]. Encapsulating antifungal drugs within nanofibers can enhance drug solubility, stability, and bioavailability, while reducing non-specific toxicity. Controlled release from nanofibers ensures sustained drug delivery to the infection site, minimizing systemic toxicity and maintaining therapeutic concentrations at the target site, potentially reducing administration frequency. Various methods are available for nanofiber production, including drawing, template synthesis, self-assembly, and electrospinning [6]. Electrospinning, the method selected this study, offers several advantages over other methods: It is relatively simple, versatile, allowing the use of various polymers and drugs, scalable for large-scale production, and cost-effective [6]. This justifies our choice of electrospinning for this research.

This study develops and evaluates a novel antifungal formulation using electrospun nanofibers loaded with polyvinyl alcohol (PVA), nystatin (NYS) and thymol (THY) (PVA-NYS-THY) to combat *C. glabrata* infections. NYS, a polyene antifungal effective against *Candida* species, targets ergosterol in fungal membranes, disrupting membrane integrity and causing cell death [7]. However, its clinical use is limited by poor solubility, instability, toxicity, and low bioavailability [8]. Encapsulation within nanofibers aims to overcome these limitations. THY, a natural phenolic monoterpene with antifungal properties, disrupts membrane integrity, binds to ergosterol, and increases membrane permeability, thereby facilitating antifungal entry [9]. Combining NYS and THY is hypothesized to produce a synergistic effect against *C. glabrata*, potentially reducing the required NYS dose and minimizing toxicity [10]. This study specifically examines the impact of PVA-NYS-THY nanofibers on the expression of key virulence genes, *SAP* and *ALS*, in *C. glabrata*. SAP enzymes facilitate tissue invasion and immune evasion, while ALS glycoproteins mediate adhesion and biofilm formation [11]. We hypothesize that PVA-NYS-THY nanofibers will modulate the expression of these genes, reducing *C. glabrata* virulence. Our goal is to assess PVA-NYS-THY nanofibers as an innovative antifungal formulation to enhance NYS efficacy against this challenging pathogen.

2. Materials and Methods

2.1. Sample collection and preparation

Oral and vaginal swabs were collected from 100 patients with suspected candidiasis within October 2023 to March 2024. The samples were cultured on *Candida* Chrome Agar and incubated at 30 ˚C for 72 hours. Yeast colonies were initially identified based on morphological and biochemical characteristics. Twelve *C. glabrata* isolates were selected for further molecular characterization and antifungal susceptibility testing.

2.2. Molecular confirmation

Genomic DNA was extracted from the 12 selected isolates using the phenol-chloroform method. The *18S rRNA* gene was amplified by polymerase chain reaction (PCR) using specific primers (F: AGCTGGTTGATTCTGCCAG, R: TGATCCTCCYGCAAGTTCAC), then purified, and sequenced (Pishgam Biotechnology Company). Sequence alignment (ClustalW) and phylogenetic analysis (MEGA7) confirmed the isolates as *C. glabrata*. PCR was also used to detect *ALS* and *SAP* gene families using specific primers.

2.3. Materials and nanofiber preparation

THY ($C_{10}H_{14}O$, 150.22 g/mol, EC 201-944-8) and NYS (Sigma-Aldrich) met analytical standards as outlined by Ph. Eur., BP, NF, and USP. Stock solutions (10 mg/mL) were prepared in dimethyl sulfoxide (DMSO) with the final DMSO concentration in all experiments was kept below 1% (v/v). PVA (89,000–98,000 g/mol) was dissolved in distilled water (10% w/v) and stirred at 80 ˚C for 4 hours. Three nanofiber formulations were prepared via electrospinning (Nanospinner NS24, Inovenso, Turkey): PVA-NYS (1% w/v), PVA-THY (1% w/v), and PVA-NYS-THY (0.5% w/v each). Electrospinning parameters were optimized as follows: flow rate (1 mL/h), voltage (15 kV), needle-collector distance (15 cm), 25 ˚C, and 40% humidity. Nanofibers were collected on aluminum foil and subsequently dried in a vacuum oven at 40 ˚C for 24 hours.

2.4. Nanofiber characterization and drug release

Nanofiber morphology (diameter) was assessed using scanning electron microscopy ([SEM], JSM-6390LV, JEOL, Japan) following gold coating. ImageJ software was used to analyze fiber diameter. Fourier-transform infrared (FTIR) spectroscopy confirmed the presence of NYS and THY within the PVA matrix. Drug release kinetics were studied using a dialysis method [12]. Nanofibers containing 10 mg of drug were placed in dialysis bags (12,000-14,000 Da cutoff) and immersed in 50 mL phosphate-buffered saline ([PBS]), pH 7.4) with 0.5% Tween 80. The bags were incubated at 37 ˚C with agitation (100 rpm). At predetermined intervals, 1 mL aliquots were sampled and replaced with fresh PBS. Released drug concentrations were quantified using high-performance liquid chromatography (HPLC).

2.5. Antifungal susceptibility testing

Antifungal activity was assessed against *Candida albicans* ATCC 10231 and clinical *C. glabrata* isolates using two methods: A modified agar diffusion assay and broth microdilution for minimum inhibitory concentration/minimum fungicidal concentrations (MIC/MFC) determination.

2.5.1. Agar diffusion assay

Fungal suspensions were prepared in Mueller-Hinton broth (MHB), incubated at 30 ˚C for 18-24 hours, and adjusted to a 0.5 McFarland standard (approximately 1-5×10^6 CFU/mL) (CLSI M27, 2022). A 1:100 dilution (approximately 1-5×10^4 CFU/mL) was used to enhance visualization of inhibition zones. Six-millimeter nanofiber discs were placed on Mueller-Hinton agar (MHA) plates inoculated with the fungal suspensions using a sterile swab to create a confluent lawn. Plates were incubated at 30 ˚C for 24-48 hours. Inhibition zone diameters were measured using a digital caliper. All tests were performed in triplicate (n=3), and results are presented as Mean±SD.

2.5.2. Broth microdilution assay (MIC/MFC)

MIC and MFC values for NYS, THY, and the PVA-NYS-THY combination were determined using the broth microdilution method, according to CLSI M27 (2022) and adapted for testing of nanofibers. Serial two-fold dilutions of each agent were prepared in 96-well microtiter plates using 3-(N-morpholino) propanesulfonic acid (MOPS)-buffered Roswell Park Memorial Institute (RPMI) 1640 medium. The tested concentrations were: NYS (0.039–125 µg/mL), THY (0.625–2000 µg/mL), and PVA (0.078–25 mg/mL). DMSO, used as a solvent for THY, was kept below 1% (v/v). A 100 µL inoculum (prepared as in section 2.5.1) was added to 100 µL of drug solution per well (200 µL final volume). Plates were incubated at 30 ˚C for 24-48 hours. MIC was determined as the lowest concentration inhibiting visible growth. To determine MFC, 10 µL from no-growth wells (at or above the MIC) was subcultured onto Sabouraud dextrose agar (SDA) and incubated at 30 ˚C for 48 hours. MFC was defined as the lowest concentration with no growth on SDA.

2.6. Fractional inhibitory concentration index calculation

The fractional inhibitory concentration index (FICI) was calculated to evaluate the interactions between the antifungal agents in the combinations. The FICI was determined by adding the FIC values of each drug in the combination, where FICI=MIC of the drug in combination/MIC of the drug alone. The FICI was interpreted as follows: ≤0.5, synergism; >0.5-4, indifference; >4, antagonism (Equation 1).

1. $$FICI=\frac{MIC_A\,combination}{MIC_A\,alone}+\frac{MIC_B\,combination}{MIC_B\,alone}$$

where MIC_A combination and MIC_B combination are the MICs of drug A and drug B when used in combination, and MIC_A alone and MIC_B alone are the MICs of drug A and drug B when used alone.

Table 1. Primers sequence employed in qRT-PCR analyses [20]

Primer	Sequence	Tm (C°)	Amplicon Size (bp)
ALS-F	CTGGACCACCAGGAAACACT	59.53	226
ALS-R	GGTGGAGCGGTGACAGTAGT	61.53	
SAP-F	ACCGTTGGATTTGGTGGTGTTT	61.21	218
SAP-R	ATTATTTGTCCCGTGGCAGCAT	60.95	
ACT-F	TTGCCACACGCTATTTTGAG	55	167
ACT-R	ACCATCTGGCAATTCGTAGG	55	

2.7. Real time PCR

Real-time PCR was used to evaluate the effect of NYS- and THY-incorporated PVA nanofibers on *SAP* and *ALS* gene expression in *C. glabrata* isolates. The isolates were treated with sub-MIC concentrations of nanofibers. Total RNA was extracted (RNeasy Mini Kit, Qiagen), and cDNA was synthesized from 1 µg RNA (RevertAid First Strand cDNA Synthesis Kit, Thermo Fisher Scientific). Primers targeting *SAP* and *ALS* [13] (primer sequences in Table 1) and the *ACT1* housekeeping gene were used. qPCR (20 µL) used SYBR Green Master Mix (Applied Biosystems), 10 pmol/µL primers, 2 µL cDNA, and nuclease-free water. Cycling conditions: 95 ˚C for 10 min; 40 cycles of 95 ˚C for 15 s, 60 ˚C for 30 s, and 72 ˚C for 30 s. Relative gene expression was calculated using the $2^{-\Delta\Delta Ct}$ method (StepOne Software, version 2.3, Applied Biosystems).

2.8. Statistical analysis

All experiments were performed in triplicate (n=3), and data are presented as Mean±SD. The normality of data distribution was assessed using the Shapiro-Wilk test. For comparisons between multiple groups, a one-way analysis of variance (ANOVA) was performed. If the ANOVA indicated statistically significant differences ($P \leq 0.05$), Tukey's honestly significant difference (HSD) post hoc test was used for pairwise comparisons to determine which specific groups differed significantly from each other. These analyses were performed using GraphPad Prism software, version 9.0 (GraphPad, San Diego, CA, USA).

3. Results

3.1. Clinical isolation, molecular characterization, and antifungal susceptibility of *C. Glabrata*

This study characterized 13 *C. glabrata* isolates (13% prevalence, 26% of all *Candida* species) from 100 clinical samples. Oral swabs were the primary source (69%), followed by vaginal swabs (23%) and urine (8%). Infected patients (mean age 45.2 years) were predominantly female (53.85%), with most reporting no tobacco or alcohol use (84.62%). ICU admission (30.77%) and central venous catheter presence (69.23%) were common risk factors, while vulvovaginal candidiasis (VVC) (23.08%), including recurrent VVC (23.08%), was the most frequent diagnosis. Molecular identification using *18S rRNA* gene sequencing confirmed *C. glabrata* in 8 of 11 suspected isolates, exhibiting a 1500 bp PCR product and clustering phylogenetically with reference strains. The remaining three isolates were identified as other *Candida* species. Virulence genes ALS and SAP were detected in all confirmed *C. glabrata* isolates. Morphological and biochemical analyses supported these findings. Importantly, all *C. glabrata* isolates exhibited fluconazole resistance but remained susceptible or semi-susceptible to amphotericin B, NYS, and ketoconazole.

3.2. Characterization of PVA-NYS-THY nanofibers

Optimized PVA nanofibers containing NYS and THY (3:7 ratios to PVA) yielded smooth, uniform, knot-free nanofibers, observed visually and confirmed by SEM (Figure 1), showing an average diameter of ~100 nm. FT-IR spectroscopy (Figure 2) revealed characteristic peaks for PVA, NYS, and THY, including O-H, N-H, C-H, C=O, C=C, C-N, and C-O groups. Crucially, peak shifts and splitting, such as the splitting of NYS's C=O stretch (1728 cm^{-1} to 1724 cm^{-1} and 1732 cm^{-1}) and the shift in PVA/THY's C-OH stretch (944 cm^{-1}), indicated

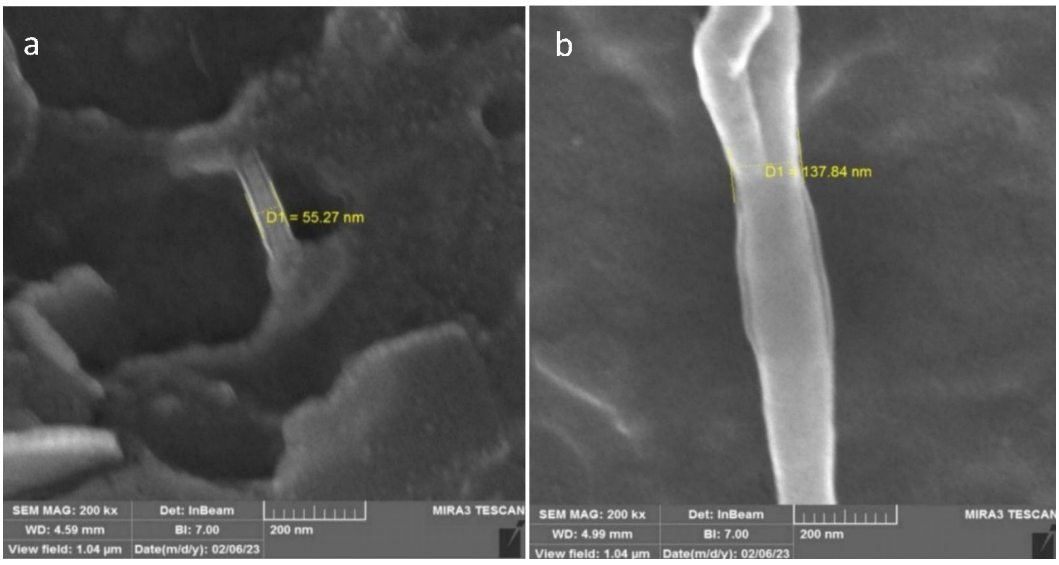

Figure 1. SEM images of PVA-NYS-THY nanofibers showing smooth and uniform morphology

A) Broader view with fiber diameter of 55.27 nm; B) Close-up view with fiber diameter of 137.84 nm.

Note: Field of view 1.04 µm, magnification k×200, sample to lens distance 4.59 mm.

intermolecular interactions, likely hydrogen bonding, within the composite nanofiber structure.

3.3. In vitro drug release profiles of PVA-based nanofibrous drug delivery systems

The in vitro drug release studies revealed distinct release kinetics profiles for the PVA-based nanofibrous systems. All formulations displayed a biphasic release pattern, characterized initially by a slow- release phase, followed by a more rapid release, culminating in a plateau. The PVA-NYS combination exhibited the most rapid and substantial release, reaching 95% of the drug payload within 76 hours (Figure 3). The PVA-THY combination demonstrated a comparable release rate, achieving 86% release within the same timeframe. In contrast, the PVA-NYS-THY triple-drug delivery system displayed a slower, more sustained release profile, reaching 84% release after 80 hours.

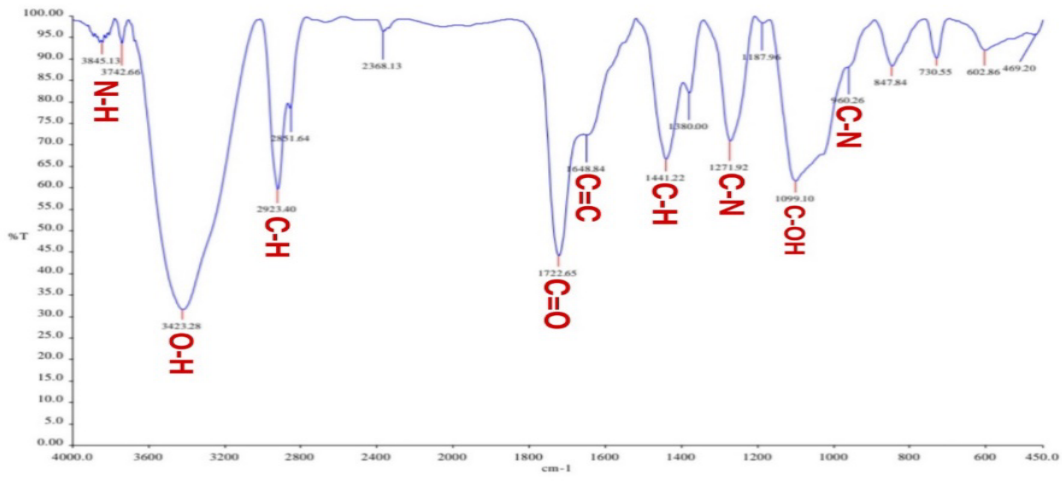

Figure 2. FT-IR spectrum of PVA-NYS-THY nanofibers

Note: The spectrum reveals the functional groups and the interactions of the compounds. The peaks of O-H, C-H, C=O, C=C, C-N, and C-O bonds are observed. The peak of C=O bond of NYS splits and the peak of C-O bond of PVA or THY shifts, indicating complexation and intermolecular forces. The interactions suggest hydrogen bonds between the hydroxyl, carbonyl, and amine groups of PVA, NYS, and THY.

Table 2. Comparison of the antibacterial activity of three different drug combinations against eight *C. glabrata* isolates

Isolates	Mean±SD		
	PVA-NYS	PVA-THY	PVA-NYS-THY
A1	30±3.1	14±0.9	35±3.4
A2	30±1.4	12±1.7	37±2.7
A3	31±1.8	13±1.1	33±0.8
A4	27±3.5	14±0.8	35±2.6
A5	30±2.7	15±1.3	34±2.5
A6	30±4.2	13±0.9	37±3.1
A7	31±1.5	14±1.2	35±2.2
A8	30±2.6	15±0.6	35±2.3

Note: The values represent the diameter of the inhibition zone (in millimeters) measured after 24 hours of incubation.

3.4. Qualitative antibiogram test results

Qualitative antibiograms revealed that PVA-based nanofibrous scaffolds exhibited antifungal activity against *C. albicans* (control) and *C. glabrata*. Inhibition zones varied across the different formulations (Table 2). Importantly, the PVA-NYS-THY composite demonstrated the most potent activity against *C. glabrata*, with inhibition zones between 33 to 37 mm. The PVA-NYS combination also demonstrated robust activity, exhibiting inhibition zones between 27 and 31 mm. In contrast, the PVA-THY scaffold displayed the lowest antifungal activity, with zones between 12 to 15 mm.

3.5. MIC, MFC, and FICI results

The antifungal potency of NYS, THY, and PVA, alone and in combination, was assessed against *C. glabrata* by determining MIC, MFC, and FIC indices (Table 3). Results indicated that PVA-based nanofibers exhibited enhanced antifungal activity compared to the individual drugs. Both PVA-NYS and PVA-THY formulations displayed lower MIC and MFC values than their respective free drug counterparts, suggesting that PVA potentiated the antifungal efficacy of NYS and THY. The FIC values for PVA-NYS and PVA-THY (both 0.25) further confirmed this synergistic effect, indicating that the combined action of the drugs with PVA exceeded the additive effect. Importantly, the PVA-NYS-THY nanofibers demonstrated even more pronounced synergy, with a lower FIC value of 0.125.

3.6. Modulation of *C. glabrata* virulence gene expression by antifungal nanofibrous systems

This study investigated the impact of antifungal agents NYS and THY, alone and in combination with PVA, on the expression of virulence genes *ALS* and *SAP* in *C. glabrata*. Both genes were significantly downregulated in all treatment groups compared to the control. NYS alone reduced *ALS* expression by 51% and *SAP* by 49%, with further reductions observed when combined with PVA.

Table 3. MIC, sub MIC, MFC, and FICI values in different drug groups against *C. glabrata*

Drug Formulation	MIC (µg/mL)	Sub MIC (µg/mL)	MFC (µg/mL)	FICI
NYS	62.5	31.25	125	-
NYS-PVA	15.62	7.81	31.25	0.25
THY	250	125	500	-
THY-PVA	62.5	31.25	125	0.25
NYS-THY	62.5	31.25	125	-
NYS-THY-PVA	7.81	3.9	15.62	0.125

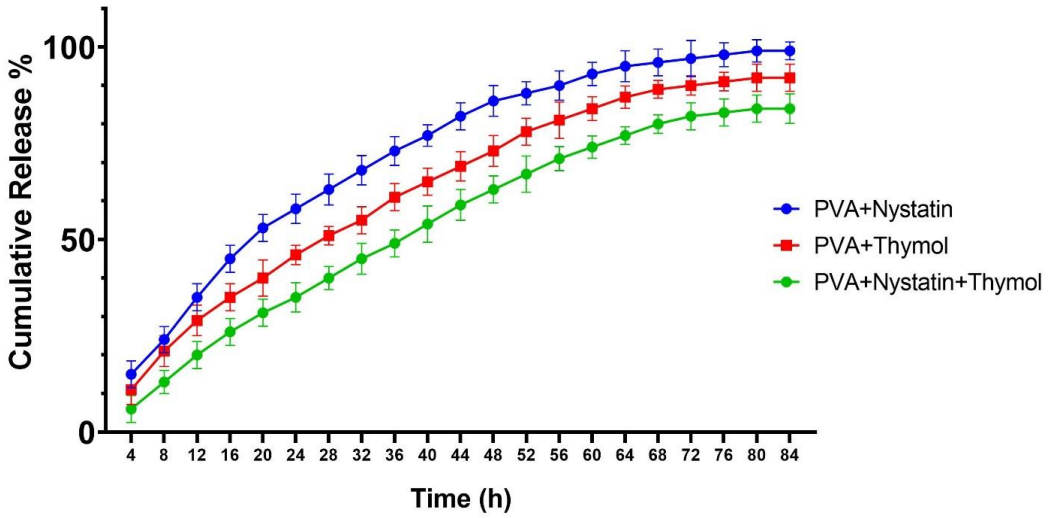

Figure 3. Drug release profiles of PVA-NYS, PVA-THY, and PVA-NYS-THY nanofibers

Note: The drug release curves showed a controlled pattern. The PVA-NYS nanofibers had the highest and fastest drug release, reaching 95% after 76 hours. The PVA-THY nanofibers had the second highest and fastest drug release, reaching 86% after 76 hours. The PVA-NYS-THY nanofibers had the lowest and slowest drug release, reaching 84% after 80 hours.

THY alone decreased *ALS* by 47% and *SAP* by 42%, also enhanced by PVA addition. The most substantial down-regulation was achieved with the combined treatment of NYS, THY, and PVA, resulting in a 65% reduction in ALS and a 68% reduction in SAP expression (Figure 4).

4. Discussion

In this study, we developed PVA-NYS-THY nanofibers to target *C. glabrata*, a fluconazole-resistant pathogen. Clinical strains of *C. glabrata* were isolated from patient specimensand characterized using various analytical techniques, and their antifungal efficacy was assessed through MIC, MFC, and FICI assays. We also investigated *ALS* and *SAP* gene expression, crucial for *C. glabrata* virulence, using PCR and qRT-PCR methods.

Our findings revealed that *C. glabrata* accounted for 13% of the total isolates and 26% of the *Candida* isolates, consistent with previous reports that identify *C. glabrata* as the second most prevalent *Candida* species after *C. albicans* [14]. The majority of the *C. glabrata* isolates were obtained from oral swabs (69%), followed by vaginal swabs (23%) and urine samples (8%), reflecting the typical distribution of candidiasis across different anatomical sites [15]. Morphological and biochemical analyses confirmed the classical characteristics of *C. glabrata*. Antifungal susceptibility testing showed that all *C. glabrata* isolates were resistant to fluconazole, consistent with the species' propensity for azole resistance, often mediated by efflux pump overexpression and *ERG11* mutations [4]. However, the isolates were susceptible or semi-susceptible to polyene antifungals, including amphotericin B and NYS, and exhibited susceptibility to ketoconazole. These findings highlight the need to explore mechanisms driving variable azole responses for better candidiasis treatment strategies.

We synthesized PVA-NYS-THY nanofibers using electrospinning, with PVA as the carrier polymer, and NYS and THY as the active antifungal agents. PVA's biodegradability and biocompatibility make it ideal for forming smooth, uniform nanofibers [16]. NYS and THY were chosen for their established antifungal properties, low molecular weights, and polarities, facilitating their integration with PVA [7]. These agents can form hydrogen bonds with PVA, potentially enhancing the nanofibers' stability and uniformity [16]. Formulation optimization ensured defect-free fibers. SEM revealed uniform nanofibers with an average diameter of ~200 nm. FT-IR spectroscopy confirmed the presence of characteristic functional groups and interactions, indicating structural integrity and functional performance of the nanofibers [17]. These results demonstrate PVA-NYS-THY nanofibers as a promising platform for antifungal drug delivery.

The controlled release behavior of PVA-NYS-THY nanofibers was monitored, revealing a pattern suitable for sustained drug delivery. The solubility and affinity of NYS and THY for the PVA matrix influenced their release profiles [18]. NYS, with higher solubility and

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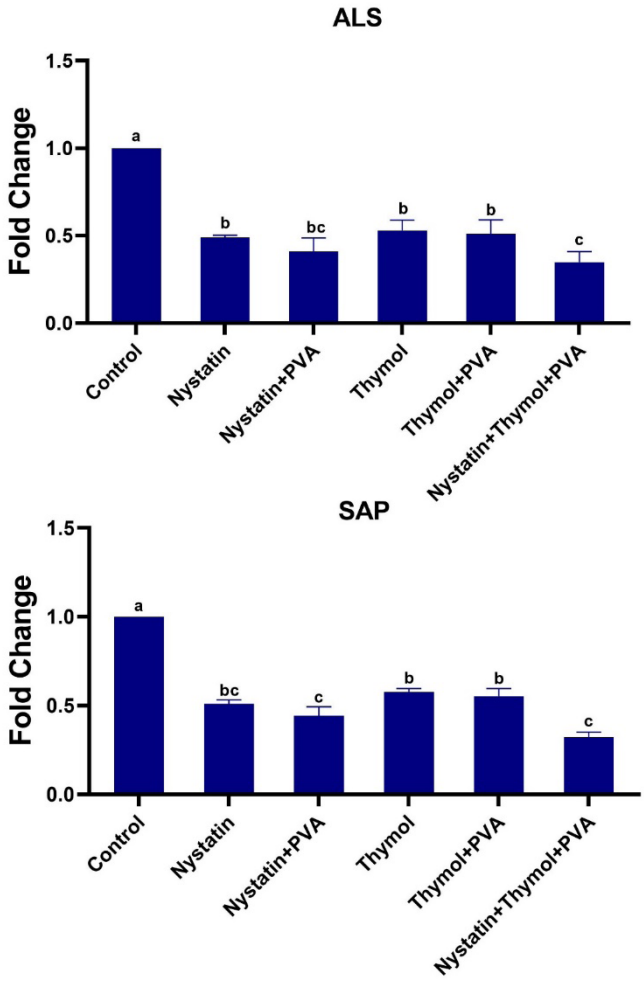

Figure 4. Expression of *ALS* and *SAP* genes in *C. glabrata* under different antifungal treatments

Note: The expression of *ALS* and *SAP* genes, which encode for ALS proteins and SAP, respectively, was measured by RT-PCR in *C. glabrata* isolates exposed to various antifungal drugs: NYS, THY, PVA, or their combinations. The expression levels were normalized to the control group, which received no treatment. The results showed that the expression of both genes decreased significantly in all drug groups compared to the control group ($P<0.05$). The control group had the highest expression level of both genes, followed by THY, THY+PVA, and NYS groups, which had similar expression levels. The NYS+PVA and NYS+THY+PVA groups had the lowest expression level of both genes. Means followed by the same letter do not differ significantly by Tukey test.

lower PVA affinity, exhibited a more rapid initial release, while THY, with moderate solubility and affinity, showed a more sustained release. The combination within the nanofiber matrix resulted in a release profile that combined aspects of both individual drugs, demonstrating the potential for controlled and prolonged release. These findings suggest the potential of PVA-NYS-THY nanofibers for customizable drug delivery.

The antifungal efficacy of PVA-NYS-THY nanofibers against *C. glabrata* was assessed through inhibition zones, MIC, and MFC measurements. The results demonstrated that antifungal activity correlated with the concentration and composition of NYS and THY, which is related to the drug release kinetics from the nanofiber matrix [19]. NYS disrupts the fungal cell membrane by binding to ergosterol, while THY's mechanism of action involves disruption of membrane integrity, binding to ergosterol, and increased membrane permeability, facilitating antifungal entry [9]. Resistance mechanisms in *C. glabrata* involve reduced ergosterol for NYS and efflux pump upregulation or enzyme modification for THY [20]. PVA-NYS-THY nanofibers exhibited the most potent activity, with the largest inhibition zones and lowest MIC and MFC values. The synergistic effect of NYS and THY, demonstrated by the FICI, enhances the therapeutic potential of this formulation against fluconazole-resistant *C. glabrata*. This synergy is likely due to the

combined membrane disruption and other mechanisms of action of the two drugs.

We investigated the presence of ALS and SAP genes in *C. glabrata* isolates using PCR. These genes encode agglutinin-like sequence (ALS) proteins and secreted aspartyl proteinases (SAP), key virulence factors involved in adhesion and pathogenicity [11]. All tested *C. glabrata* isolates (n=8) were positive for both genes, suggesting their consistent co-occurrence and potential role in facilitating invasive infections [21]. The co-expression of *ALS* and *SAP* may enhance the organism's ability to colonize host tissues and evade immune defenses, making them important targets for future therapeutic strategies. Further research should investigate the regulation of these genes and their precise roles in antifungal resistance and virulence.

We investigated the effects of NYS and THY, alone and in combination with PVA, on *ALS* and *SAP* gene expression in *C. glabrata* [22]. NYS alone reduced *ALS* and *SAP* expression, while THY also decreased the expression of both genes. The combination of NYS and THY with PVA further amplified these effects, with the PVA-NYS-THY combination showing the greatest reduction in both *ALS* and *SAP* expression. This enhanced effect likely results from the combined mechanisms of action of NYS and THY (membrane disruption and oxidative stress, respectively) [20], potentially enhanced by PVA's influence on drug stability and delivery. This downregulation of virulence genes suggests a potential mechanism by which the nanofibers exert their antifungal effect. The downregulation of *ALS* and *SAP* may be linked to disruption of regulatory pathways involving transcriptional regulators like Rim101p, Hap43p, and Efg1p [23]. NYS and THY may influence these regulators through mechanisms such as altering ambient pH (activating Rim101p, repressing ALS) or generating reactive oxygen species (ROS) (activating Hap43p, repressing *SAP*) [24]. Disruption of ergosterol biosynthesis can also affect regulators like Efg1p [25]. Further studies are needed to elucidate the precise molecular mechanisms involved. The enhanced downregulation of *ALS* and *SAP* in the PVA-NYS-THY group suggests that combining these agents within the nanofiber formulation can effectively reduce *C. glabrata* virulence.

This study presents significant implications for future research and clinical practice. It demonstrates electrospinning as a promising technique for developing antifungal formulations that deliver natural agents like NYS and THY in a controlled manner. It also shows that NYS and THY act synergistically against *C. glabrata*, enhancing their antifungal activity. Furthermore, the study reveals that these agents can modulate *ALS* and *SAP* gene expression, potentially reducing *C. glabrata* virulence. Future research should assess biocompatibility and conduct in vivo testing to evaluate the clinical potential of these nanofibers.

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Compliance with ethical guidelines

This study was approved by the Research Ethics Committees of North Tehran Branch, Islamic Azad University (IAU), Tehran, Iran, and certify that this study was performed in accordance with the ethical standards as laid down in the 1964 Declaration of Helsinki and its later amendments or comparable ethical standards.

Data availability

The data supporting the findings of this study are available on request from the corresponding author upon reasonable request.

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Authors' contributions

Conceptualization, study design, review and editing: Mohaddeseh Larypoor and Mansour Bayat; Data analysis and interpretation: Romina Ahmadian and Mohaddeseh Larypoor; Statistical analysis: Romina Ahmadian1 and Mohaddeseh Larypoor; Writing the original draft: Romina Ahmadian.

Conflict of interest

The authors declared no conflict of interest.

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